## SEQUENCE LISTING

- <110> Benson, Timothy E
- <120> CRYSTALLIZATION AND STRUCTURE DETERMINATION OF STAPHYLOCOCCUS AUREUS THIOREDOXIN REDUCTASE
- <130> 00032.US1
- <140> Unassigned
- <141> 2001-04-03
- <150> 60/195,055
- <151> 2000-04-06
- <160> 3
- <170> PatentIn Ver. 2.1
- <210> 1
- <211> 320
- <212> PRT
- <213> Staphylococcus aureus
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- Glu Val Glu Asn Phe Pro Gly Phe Glu Met Ile Thr Gly Pro Asp Leu
  50 55 60
- Ser Thr Lys Met Phe Glu His Ala Lys Lys Phe Gly Ala Val Tyr Gln 65 70 75 80
- Tyr Gly Asp Ile Lys Ser Val Glu Asp Lys Gly Glu Tyr Lys Val Ile
  85 90 95
- Asn Phe Gly Asn Lys Glu Leu Thr Ala Lys Ala Val Ile Ile Ala Thr 100 105 110
- Gly Ala Glu Tyr Lys Lys Ile Gly Val Pro Gly Glu Gln Glu Leu Gly
  115 120 125

Gly Arg Gly Val Ser Tyr Cys Ala Val Cys Asp Gly Ala Phe Phe Lys 130 135 140

Gly Thr Phe Thr Thr Lys Phe Ala Asp Lys Val Thr Ile Val His Arg 165 170 175

Arg Asp Glu Leu Arg Ala Gln Arg Ile Leu Gln Asp Arg Ala Phe Lys 180 185 190

Asn Asp Lys Ile Asp Phe Ile Trp Ser His Thr Thr Lys Ser Ile Asn 195 200 205

Glu Lys Asp Gly Lys Val Gly Ser Val Thr Leu Thr Ser Thr Lys Asp 210 215 220

Gly Ser Glu Glu Thr His Glu Ala Asp Gly Val Phe Ile Tyr Ile Gly 225 230 235 240

Met Lys Pro Leu Thr Ala Pro Phe Lys Asp Leu Gly Ile Thr Asn Asp 245 250 255

Val Gly Tyr Ile Val Thr Lys Asp Asp Met Thr Thr Ser Val Pro Gly 260 265 270

Ile Phe Ala Ala Gly Asp Val Arg Asp Lys Gly Leu Arg Gln Ile Val 275 280 285

Thr Ala Thr Gly Asp Gly Ser Ile Ala Ala Gln Ser Ala Ala Glu Tyr 290 295 300

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<211> 320

<212> PRT

<213> Escherichia coli

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Leu	Ile	Thr 35	Gly	Met	Glu	Lys	Gly 40	Gly	Gln	Leu	Thr	Thr 45	Thr	Thr	Glu
Val	Glu 50	Asn	Trp	Pro	Gly	Asp 55	Pro	Asn	Asp	Leu	Thr 60	Gly	Pro	Leu	Leu
Met 65	Glu	Arg	Met	His	Glu 70	His	Ala	Thr	Lys	Phe 75	Glu	Thr	Glu	Ile	Ile 80
Phe	Asp	His	Ile	Asn 85	Lys	Val	Asp	Leu	Gln 90	Asn	Arg	Pro	Phe	Arg 95	Leu
Asn	Gly	Asp	Asn 100	Gly	Glu	Tyr	Thr	Cys 105	Asp	Ala	Leu	Ile	Ile 110	Ala	Thr
Gly	Ala	Ser 115	Ala	Arg	Tyr	Leu	Gly 120	Leu	Pro	Ser	Glu	Glu 125	Ala	Phe	Lys
Gly	Arg 130	Gly	Val	Ser	Ala	Cys 135	Ala	Thr	Cys	Asp	Gly 140	Phe	Phe	Tyr	Arg
Asn 145	Gln	Lys	Val	Ala	Val 150	Ile	Gly	Gly	Gly	Asn 155	Thr	Ala	Val	Glu	Glu 160
Ala	Leu	Tyr	Leu	Ser 165	Asn	Ile	Ala	Ser	Glu 170	Val	His	Leu	Ile	His 175	Arg
Arg	Asp	Gly	Phe 180		Ala	Glu	Lys	Ile 185	Leu	Ile	Lys	Arg	Leu 190	Met	Asp
Lys	Val	Glu 195		Gly	Asn	Ile	Ile 200	Leu	His	Thr	Asn	Arg 205		Thr	Glu
Glu	Val 210	Thr	Gly	Asp	Gln	Met 215		Val	Thr	Gly	Val 220	Arg	Leu	Arg	Asp
Thr 225	Gln	Asn	Ser	Asp	Asn 230	Ile	Glu	Ser	Leu	Asp 235		Ala	Gly	Leu	Phe 240
Val	Ala	Ile	Gly	His 245		Pro	Asn	Thr	Ala 250		Phe	Glu	Gly	Gln 255	Leu

Glu Leu Glu Asn Gly Tyr Ile Lys Val Gln Ser Gly Ile His Gly Asn

260 265 270

Ala Thr Gln Thr Ser Ile Pro Gly Val Phe Ala Ala Gly Asp Val Met 275 280 285

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Ala Ala Leu Asp Ala Glu Arg Tyr Leu Asp Gly Leu Ala Asp Ala Lys 305 310 315 320

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<212> PRT

<213> Arabidopsis thaliana

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Lys Pro Leu Leu Phe Glu Gly Trp Met Ala Asn Asp Ile Ala Pro Gly 35 40 45

Gly Gln Leu Thr Thr Thr Thr Asp Val Glu Asn Phe Pro Gly Phe Pro 50 55 60

Glu Gly Ile Leu Gly Val Glu Leu Thr Asp Lys Phe Arg Lys Gln Ser 65 70 75 80

Glu Arg Phe Gly Thr Thr Ile Phe Thr Glu Thr Val Thr Lys Val Asp
85 90 95

Phe Ser Ser Lys Pro Phe Lys Leu Phe Thr Asp Ser Lys Ala Ile Leu 100 105 110

Ala Asp Ala Val Ile Leu Ala Ile Gly Ala Val Ala Lys Arg Leu Ser 115 120 125

Phe Val Gly Ser Gly Glu Val Leu Gly Gly Phe Trp Asn Arg Gly Ile 130 135 140 Pro Leu Ala Val Ile Gly Gly Gly Asp Ser Ala Met Glu Glu Ala Asn 165 170 175

Phe Leu Thr Lys Tyr Gly Ser Lys Val Tyr Ile Ile His Arg Arg Asp 180 185 190

Ala Phe Arg Ala Ser Lys Ile Met Gln Gln Arg Ala Leu Ser Asn Pro 195 200 205

Lys Ile Asp Val Ile Trp Asn Ser Ser Val Val Glu Ala Tyr Gly Asp 210 215 220

Gly Glu Arg Asp Val Leu Gly Gly Leu Lys Val Lys Asn Val Val Thr 225 230 235 235

Gly Asp Val Ser Asp Leu Lys Val Ser Gly Leu Phe Phe Ala Ile Gly 245 250 255

His Glu Pro Ala Thr Lys Phe Leu Asp Gly Gly Val Glu Leu Asp Ser 260 265 270

Asp Gly Tyr Val Val Thr Lys Pro Gly Thr Thr Gln Thr Ser Val Pro 275 280 285

Gly Val Phe Ala Ala Gly Asp Val Gln Asp Lys Lys Tyr Arg Gln Ala 290 295 300

Ile Thr Ala Ala Gly Thr Gly Cys Met Ala Ala Leu Asp Ala Glu His 305 310 310 315 320

Tyr Leu Gln Glu Ile Gly Ser Gln Glu Gly Lys Ser Asp 325 330